## RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:

Source:

Date Processed by STIC:

ENTERED



IFW16

RAW SEQUENCE LISTING DATE: 01/04/2005
PATENT APPLICATION: US/09/312,596F TIME: 13:26:37

Input Set : A:\593601.txt

Output Set: N:\CRF4\01042005\I312596F.raw

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3 <110> APPLICANT: Role, Lorna W.
         Talmage, David
         Bao, Jianxin
 7 <120> TITLE OF INVENTION: A-FORM OF CYTOPLASMIC DOMAIN OF nARIA (CRD-NEUREGULIN
         AND USES THEREOF
10 <130> FILE REFERENCE: 0575/59360
12 <140> CURRENT APPLICATION NUMBER: 09/312,596F
13 <141> CURRENT FILING DATE: 1999-05-14
15 <160> NUMBER OF SEQ ID NOS: 5
17 <170> SOFTWARE: PatentIn Ver. 2.1
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 3212
21 <212> TYPE: DNA
22 <213> ORGANISM: CHICKEN nARIA
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27 aaggtgagag tettgeettt egetetggge tattggttea ettaateegg teaatttgtt 180
28 cgctgctcgt ggttgtcttt ctccccgccc tccttccccc tgttttgttt tgtttcgctt 240 ·
29 gctttcgggg ggacgctcct tccctcagtc agaagagctg gaattgcttg agaggcgtat 300
30 aaggaattat aaaagtggcc aggaaacacg agcgcagtga ctgcagagct gcccttggct 360
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40 gtatgactct cctacacacc ttgaccctgg gaggatagga caagacccaa ggagcactgt 960
41 ggatcctaca gctctgtctg cctgggtgcc ttcggaggtg tatgcctcac ccttccccat 1020
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43 caggeeette etteageett etetetaeaa eegeateeta gatgtegggt tgtggteete 1140
44 tgccacaccg tcactgtcac catcctccct ggagcctacc acggcatctc aggcacaagc 1200
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52 catggcaaat gggccacacc accccaaccc accaccagac aatgtccagc tggtgaatca 1680
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54 gttttccaca agccactaca cctcaacaac tcatcactcc atgacagtca cccagacgcc 1800
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56 cgtcagctcc tcagtggaga atagcaggca caccagccca acagggccac gaggccgcct 1920
57 caatggcatt ggtgggccaa gggaaggcaa cagcttcctc cggcatgcaa gagagacccc 1980
58 tgactcctac cgagactctc ctcacagtga aaggtatgtc tcagctatga ccacaccagc 2040
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61 ggacgaggag agaccgctgc tgttggtgac cccaccacgg ctgcgtgaga agtacgacaa 2220
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71 ttccgctgat agaaaacaag agtggaaaaa gaaactttta taaattaagt atacgtatgt 2820
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73 tatcaatggt gcctttatgt tatgttatgt tgagagcaag ttttgtacag ctacaatgat 2940
74 tgctgtcccg tagtattttg caaaaccttc tagccctcag ttgttctggc ttttttgtgc 3000
75 attgcattat aatgactgga tgtatgattt gcaagaattg cagaagtccc catttgcttg 3060
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PATENT APPLICATION: US/09/312,596F TIME: 13:26:37

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176 Glu Thr Phe Pro Ser Pro Ser Ala Gln Leu Ser Pro Asp Ala Ser Leu

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177		210					215					220				
	Gly												His	Ara	Glu	Asp
	225	_	LÇU		1114	230	014				235		1110	**** 9	02.4	240
	Ser		Val	Pro	Glv							Thr	Cvs	Cvs	Val	
183		*****	VUI		-		1124	_		250	DCI	1111	Cys	Cys	255	Cys
	Leu	Glu	Δla	Glu							Δsn	Ser	Glu	T.vs		Cvs
	пец			260	nry	neu	цуз	Gry	_		ASII	261	GIU	270	110	Cys
	Ile				Leu	Δla	C178	T.A11				Cve	T.611		Tla	Δla
189		ATa	275	116	пец	ALG	Cys	280				Cys	285	Суз	116	ATG
	Gly	Τ.Δ11		ጥፖኮ	V = 1	Dha	Val					Glu		Aen	Sar	Pro
192	_	290	цуэ	тър	Val	1116		АЗР	_			300	т <b>у т</b>	АЗР	561	110
	Thr		T.A11	Asn	Pro	Glv							Δra	Sar	Thr	Val
	305	1115	пси	ASP	110	_	nrg		_	OTII	_	110	-	561	1111	320
	Asp	Pro	Thr	Δla	T.e.11					Pro				ጥህዮ	Δla	
198	_	110	1111	7114	325	501	71 <u>1</u> C	111	٧۵٢	330	DCI	OIU	Val	1 7 1.	335	DCI
	Pro	Phe	Pro	Tle		Ser	T.e.ii	Glu	Ser		Δla	Glu	Val	Thr		Gln
201		1110	110	340			пси			_	ΛIα		Val	350	٧۵٢	0111
	Thr	Asp	Ser										Gln		Ser	T.e.11
204		7100	355		ПСС		110		**** 9			БСС	365	110	501	DC u
	Tyr	Asn					Val					Ser		Thr	Pro	Ser
207		370		110			375		шси			380	11,1,0	1111		
	Leu												Gln	Ala	Gln	Ala
	385	001		<b>-</b>	<b>001</b>	390					395	001	01		<b></b>	400
	Thr	Glu	Thr	Asn	Leu							Ser	Thr	Ser	Thr	
213		014			405					410	200			001	415	J J _
	Thr	Thr	Glv	Thr							Asp	Ile	Lvs	Gln	_	Ala
216			<u>1</u>	420					425	-	-		_	430	-1-	
	Phe	Cvs	Val												Pro	Asn
			435		_	_		440				1	445			
	Pro			Tyr	Leu	Cvs	Arq	Cvs				Phe	Thr	Glv	Asp	Arg
222		450	_	-		_	455	_				460		-	•	,
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225	465			_		470				-	475			-		480
227	Phe	Met	Glu	Ala	Glu	Glu	Leu	Tyr	Gln	Lys	Arg	Val	Leu	Thr	Ile	Thr
228					485					490					495	
230	Gly	Ile	Cys	Ile	Ala	Leu	Leu	Val	Val	Gly	Ile	Met	Cys	Val	Val	Ala
231				500					505					510		
233	Tyr	Cys	Lys	Thr	Lys	Lys	Gln	Arg	Lys	Lys	Leu	His	Asp	Arg	Leu	Arg
234			515					520					525			
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237		530					535					540				
239	Pro	His	His	Pro	Asn	Pro	Pro	Pro	Asp	Asn	Val	Gln	Leu	Val	Asn	Gln
240	545					550					555					560
242	Tyr	Val	Ser	Lys	Asn	Ile	Ile	Ser	Ser	Glu	Arg	Val	Val	Glu	Arg	Glu
243					565					570					575	
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246				580					585					590		
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249			595					600					605			

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251 252		Glu 610		Ile			Glu 615		His			Leu 620	Val	Ser	Ser	Ser
254		Glu				His	Thr		Pro	Thr	Gly	Pro	Arg	-	_	Leu 640
			Tle	Glv	Glv										His	
		_		Ory	_		_	014	_		501			_	655	1114
260	Arg	Glu	Thr	Pro											Arg	Tyr
261				660					665					670		
									_					Asp	Phe	His
		<b>5</b>				-			5				685	75	5	** . 7
267		690		ser		_		Pro				700	Ser	Pro	Pro	Val
													Ser	Pro	Phe	Met
								0					002			720
272	Asp	Glu	Glu	Arg	Pro						Pro	Pro	Arg	Leu	Arg	Glu
273					725					730					735	
	_	_	_												Asn	Pro
				740							Ď			750	17a l	C1
		HIS		ser					Pro				765	тте	vaı	Glu
													_	Gln	Glu	Pro
282	1-0 <sub>F</sub>	770		- 1 -			775	02		- 1 -		780				
284	Pro	Lys	Lys	Leu	Thr	Asn	Ser	Arg	Arg	Val	Lys	Arg	Thr	Lys	Pro	Asn
	785					790					795					800
	Gly	His	Ile	Ser		Arg	Val	Glu	Val	-	Ser	Asp	Thr	Ser	Ser	Gln
288	Com	mb ~	C 0 ~	C 0 70	805	C 0 7	C1	mb ~	C1	810	C1	7 ~~~	Tla	C1	815	7.00
290	ser	Thr	ser	820	Glu	Ser	GIU	Thr	825	Asp	GIU	Arg	тте	830	Glu	Asp
	Thr	Pro	Phe		Ser	Ile	Gln	Asn		Met	Ala	Thr	Ser		Glu	Pro
294			835					840					845			
296	Ala	Ala	Ala	Tyr	Arg	Leu	Ala	Glu	Asn	Arg	Thr	Asn	Pro	Ala	Asn	Arg
297		850					855					860				
		Ser	Thr	Pro	Glu		Leu	Gln	Ala	Arg		Ser	Ser	Val	Ile	
	865	Cln	7 an	Dro	T10	870	W-1	Vaa	7 cn	т1.	875	Tvc	Thγ	п;с	7 ~~	880 Bho
303	ASII	GTII	ASP	FIO	885	Ala	vai	naa	ASP	890	ASII	гу	1111	птэ	Arg 895	riie
	Thr	Cys	Lys	Thr		Phe	Tyr	Ile	Met		Tyr	Ser	Thr	Phe		Leu
306		4	-	900					905	4				910	_	
308	Asn	Asn	Leu	Phe	Tyr	Phe	Ser	Asn	Ser	Ala	Asp	Arg	Lys	Gln	Glu	Trp
309			915				_	920					925			
	Lys	_	Lys	Leu	Leu	Xaa		Lys	Tyr	Thr	Tyr		Gln	Met	Cys	Tyr
312	Wa 1	930 Pro	Ф	Wal	7.1.	T10	935	Ф	202	Tla	202	940	Mot	C1,,	Tuc	7 00
	945	PIO	туг	vaı	Ald	950	rne	ıyı	ser	тте	955	гуу	Met	GTÅ	гуу	Asp 960
		Asn	Glv	Ala	Phe		Leu	Cvs	Tvr	Val		Ser	Lvs	Phe	Cys	
318	- <del>-</del>		- 1	. — •	965		<del>-</del>	1 -	1 -	970	- <del></del>		1 -		975	<del></del>
320	Ala	Thr	Met	Ile	Ala	Val	Pro	Xaa	Tyr	Phe	Ala	Lys	Pro	Ser	Ser	Pro
321			_	980				_	985					990		
323	Gln	Leu	Phe	Trp	Leu	Phe	Cys	Ala	Leu	His	Tyr	Asn	Asp	Trp	Met	Tyr

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## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:2; Xaa Pos. 32,42,113,163,182,199,888,934,984,1067 Seq#:4; Xaa Pos. 16,338,347,362,377,383,387,414,423,442

## Invalid Line Length/:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:5; Line(s) 520

VERIFICATION SUMMARY

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L:140 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:16

M:341 Repeated in SeqNo=2

L:426 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0

M:341 Repeated in SeqNo=4